



KAROLINSKA INSTITUTET
a medical university

EXHIBIT 1

Department of CLINICAL IMMUNOLOGY

Laboratory Journal No.

Name: ~~XXXXXXXXXX~~
JANS + LIND

Group:

Date: [REDACTED] from to



cont. from page no.

Study

cont. on page no.

Project no.

Study no.

27

Assembly of a synthetic gene coding for
Fcd 1 chain 1 using Tag

PCR

Oligos 132, 133, 6T, 155 10 μ M

1 μ l of each oligo 132-133

1 μ l of NTP 10 mM

0.5 μ l Tag

1 μ l 10x Tag-buffer

35 μ l H₂O

10 μ l

→ PCR Eppendorf program HAN51

94°C 1 min

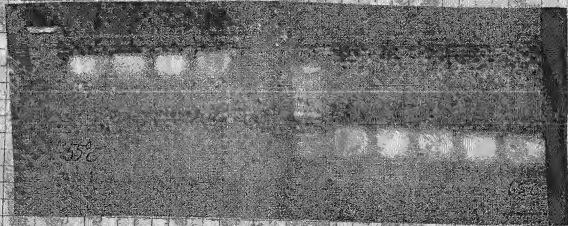
annealing 63-65°C 15 min (grad)

elongation 68°C 2 min

30 cycles ending by 10 min elongation

74°C

Result: A strong band around 300 (exp. ~ 200)



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Laboratory Journal

248910

event final page no.

Study

Assembly of chain 2. Rd. di.

control page no.

Project no.

Study no.

primers 127-131 + 138

Using 3 different DNA polymerases

Tag
ppa

Method: assemble chain 1st

Ampl. tag

expected band
at 394



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cont. from page no.

Study

cont. on page no.

Project no.

Study no.

Expression and purification of Fcd1 chain 1 and chain 2.

Fcd1 chain 1 (clone 42) and Fcd1 chain 2 (clone 29) was ligated into pET 20b and electroporated into BL-21 DE3 plys after having been cut from pTZ-Blue containing the correct sequence (see binder HCL Fcd1). Sequencing of pET 42/pET 29 was done according to standard protocol (APBT) and the result can be seen on the opposite note.

Both Fcd1 chain 1 (Fcd1:1) and Fcd1 chain 2 (Fcd1:2) was expressed according to standard protocol and purified on a HiTrap (chelex) column loaded with NiSO_4 .

Chain 1 was soluble, after ultra sonication chain 1 was found in 20mM Tris-HCl pH fraction, while chain 2 was found in the inclusion bodies after "washing" with 2M urea buffer + 20mM Tris-HCl pH 8.0. The inclusion bodies were solubilized in 6M Guanidine, transferred to 6M Urea buffer (20mM Tris-HCl pH 8.0 + 0.5M NaCl) via 6th HiTrap. Purification was done on HPLC.

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Entscheidungen und Algorithmen

[illegible]

2314

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[illegible][illegible]

20	21	22	23	24	25	26	27
0.47	11.4	265	265	2145			83

[illegible]

Grasshopper: 40 mg/m²
B = 97 mg/m²
C = 28 mg/m²
D = 66 mg/m²



cont. from page no.

cont. to page no.

Study

Project no.

Study no.

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ELISA conditions

Coating of Ed.C1 and Ed.C2 ^(100 μ l/well) over the weekend in $+4^{\circ}\text{C}$. Serum 4 times with "Tris coating". Patient's serum was added, 100 μ l/well and incubated at $+4^{\circ}\text{C}$ o.n. Wash 4 times (Wallac ELISA-washer) with "Tris coating" and 100 μ l Rabbit human anti-glyc det. 1500 times in "Van buffer". Incubation 2 hours in RT on shaker. Wash 4x "Tris coating" and add 100 μ l/well of Goat anti-rabbit-ALP conjugated (DAKO) for 1h. Wash 4 times and add substrate 3 tablets/15 μ l of Van buffer.

The result was read in ELISA reader after 45 min at 405 ~~nm~~ nm.

Result: 2.5 μ g/well serum to be an adequate coating concentration for both chains and chain 2. Mixing of the two chains can be done with coating concentrations 2.5 + 2.5 μ g/well.

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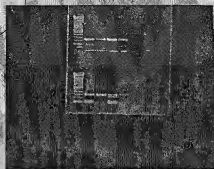
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001114

12 16 20 26

↓ ↓ ↓



1 2a 3b 4a 4b

Putting chain 1 and 2 (1+2)
together (001121)

template 1a 6ul
template clone 2g
1:10, 1:100, 1:1000

1ul
0.5ul primer 176
2ul 183
2ul dNTP
3ul 10x buff.
6ul pfu
14.5ul H₂O

30ul



↑ ↑ ↑
1:10 1:100 1:1000

Result: One band
at ~500bp which
could be clone 1+2

The bands are cut
out and purified
on QIAGEN

ligated with "perfectly
cloning kit". 10 colonies
are picked for miniprep
and possibly sequencing.

(1a) 1ul template clone 11.42 (1:1000)

2ul primer 176
2ul 183
2ul 10x buff
1ul pfu
2ul dNTP (100uM)
10ul H₂O

20ul

(1b) Some overhang
primer Tag polymerase

(2a) 1ul templ. (1:1000) clone 2g
1ul templ. 183
2ul primers 176, 183, 185
1 overhang from (1a)

(2b) Some (2a) new Tag

(3a) 1ul template clone 2 (1:1000)
2ul 183
2ul 185
1 overhang from 1a

(3b) Some (3a) new Tag

(4a) 1ul templ. 1
1ul templ. 2
2ul primer 180
1 overhang from (3a)

(4b) Some (4a) new Tag

AmpliTaq Gold
250 Units, 50ul
Store at -20°C

Applied Biosystems
A05912



cont. from page no.

Study

Linking of (chain 1 and chain 2 (scamless))
with PCR. Del d1 and Del d2

cont. on page no.

Project no.

Study no.

Aim. The aim of this experiment is to join the two chains of Fel d1 into one construct by PCR

Study outline The two sequenced chains of the major ~~chain~~ ^{allergen} of cat Fel d1 (chain 1, clone 42) and chain 2, clone 29) is joined with PCR in two steps as outlined below. In



Result (see opposite side)

Good bands of expected size was seen for both chain 1 and chain 2. But (2a) + (2b) as well as (1a) and (1b) did not work. I will continue by adding (1a) to ~~template~~ ^{template} 29 and do PCR. (Chain 1+2)

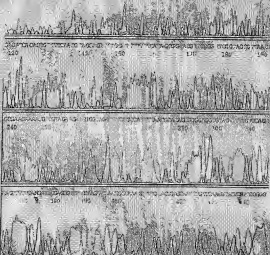


PRISM

APR 1971

F-20

APR 1971



Tel (24) 27 dfr 00105 (67-1000)

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ABI PRISM

Model 371

Version 3.4

Firmware 3.2

Line 37

Signal G112 A133 CH11 T136

OT 01101 Any Power

Seq 01101 T136

Power 1336 to 1360 Ph 11.0C 1360

Make a copy of

this file

Specify

250 260 270 280 290 300 310 320 330

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cont. on page no.

Study

Project no.

Study no.

Ligation of Pcl 1, clone 1 (2cl) and clone 5 (1cl)
into pET 20b⁺ and electroporation into BL21 plys

The fragments from Pcl 1 (2cl) clone 1 and clone 5 (1cl)
were digested purified from 1% agarose gel (2x8931)
A cleaned (Nde/Xho) pET 20b⁺ vector was used
to ligate the fragments

Conculation

(2 μ l vector fragments)
10 μ l vector
1.5 μ l 10x ATP
1.8 μ l 10x ligation buffer
1.7 μ l 1% ligase
1.3 μ l

Ligate +16°C o.n.

The ligate mix was electroporated into 50 μ l BL21-
- plys electrocompetent cells. 1 μ l ligate mix
was added to thawed cells (on ice). Electroporation
according to standard protocol. Growth on SOC
medium for 60' 37°C shaker (300 rpm) and
plated on Amp/CAN plates. One colony from
each plate was picked and grown on LB Amp
can medium, miniprep (Q-gene)
and 25 μ l of the (50 μ l) prep was
cut with Nde and Xho. Result ~ 500bp
Both clones contain the insert!!

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cont. on page no.

Purification of Fd 1 (1+2) clone 5 and
Fd 1 (2+1) clone 1 over Ni^{2+} chelate Hi-Trap

Project no.

Study no.

(in PET 200 in B2-2, P140)

1 liter of Fd 1 (1+2) and (2+1) resp. 6 medium (CAM, Amp)
was grown to 0.6 (600nm) and induced with
0.4 mM IPTG. (see 248932)

Purification according to protocol. Both proteins
were expressed as inclusion bodies and
purified accordingly. Purification on FPLC
is follow. After adsorption onto Hi-column in
6M Gna and wash also with 6 M Gna. The
column is stuck to FPLC

Program:

D conc % 0.0

A = 6 M Urea

0 ml/min 3.0 ml/min

B = 20 mM Imidazole

0 0.25 ml/min

C = 500 mM Imidazole

0 port Set 6

20 conc % B 0

80 conc % B 1.00

100 conc % B 1.00

125 conc % B 0

125 port set 6.0

MW

100

50

25

12.5

6.25

3.125

1.5625

0.78125

0.390625

0.1953125

0.09765625

0.048828125

0.0244140625

0.01220703125

0.006103515625

0.0030517578125

0.00152587890625

0.000762939453125

0.0003814697265625

0.00019073486328125

0.000095367431640625

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